

SEQUENCE LISTING

	Wolfraim, Lawrence A Letterio, John J	
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gat	t tc	c tg	g cg	c ta	c ct	c ago	c aa	c cg	g ct	g cto	g gc	c ccc	agt	t gad	c tca	576

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Cys L	ys :	Thr 35	Ile	Asp	Met	Glu	Leu 40	Val	Lys	Arg	Lys	Arg 45	Ile	Glu	Ala
Ile A	rg (Gly	Gln	Ile	Leu	Ser 55	Lys	Leu	Arg	Leu	Ala 60	Ser	Pro	Pro	Ser
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Tyr A	\sn	Ser	Thr	Arg 85	Asp	Arg	Val	Ala	Gly 90	Glu	Ser	Val	Glu	Pro 95	Glu
Pro (Glu	Pro	Glu 100		Asp	Tyr	Tyr	Ala 105	Lys	Glu	Val	Thr	Arg 110	Val	Leu
Met '	Val	Glu 115		Gly	Asn	Gln	11e	e Tyr)	: Asp	Lys	Phe	Lys 125	Gly	Thr	Pro
	Ser 130	Leu	і Туг	. Met	: Lev	ı Ph∈ 135	e Asr	n Thr	s Ser	Glu	Leu 140	Arg	Glu	Ala	Val
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			. 4		~ ~~	+ ~~	·	ים כנ	ים מי	ממ מ	cc (gga	cto	tco	acc	96

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Ile Arg Gly Gln Ile 50	e Leu Ser Lys Leu Arg 55		Pro Ser

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Met	Val	Glu 115	Ser	Gly	Asn	Gln	Ile 120	Tyr	Asp	Lys	Phe	Lys 125	Gly	Thr	Pro
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Ser	Gly	Arg	Arg	Gly 245	Asp	Leu	Ala	Thr	Ile 250	His	Gly	Met	Asn	Arg 255	Pro
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gag cag Glu Gln 95														337	
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atc Ile	ccg Pro 240	aat Asn	aaa Lys	agc Ser	gaa Glu	gag Glu 245	ctc Leu	gag Glu	gcg Ala	aga Arg	ttt Phe 250	gca Ala	ggt Gly	att Ile	gat Asp	769
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tgc Cys 335	ctt Leu	cgc Arg	cct Pro	ctt Leu	tac Tyr 340	att Ile	gat Asp	ttt Phe	aag Lys	agg Arg 345	gat Asp	ctt Leu	gga Gly	tgg Trp	aaa Lys 350	1057

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Pro 65	Glu	Val	Ile	Ser	Ile 70	Tyr	Asn	Ser	Thr	Arg 75	Asp	Leu	Leu	Gln	Glu 80
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Ile	Val 130	Arg	Phe	Asp	Val	Ser 135	Thr	Met	Glu	Lys	Asn 140	Ala	Ser	Asn	Leu
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Glu	Trp 210	Leu	His	His	Lys	Asp 215	Arg	Asn	Leu	Gly	Phe 220	Lys	Ile	Ser	Leu
His 225	Cys	Pro	Cys	Cys	Thr 230	Phe	Val	Pro	Ser	Asn 235	Asn	Tyr	Ile	Ile	Pro 240
Asn	Lys	Ser	Glu	Glu 245	Leu	Glu	Ala	Arg	Phe 250	Ala	Gly	Ile	Asp	Gly 255	Thr

Ser Thr Tyr Ala Ser Gly Asp Gln Lys Thr Ile Lys Ser Thr Arg Lys 260 Lys Thr Ser Gly Lys Thr Pro His Leu Leu Leu Met Leu Leu Pro Ser 280 Tyr Arg Leu Glu Ser Gln Gln Ser Ser Arg Arg Lys Lys Arg Ala Leu Asp Ala Ala Ser Tyr Pro Tyr Asp Val Pro Asp Tyr Ala Ser Leu Ala 310 315 Leu Asp Ala Ala Tyr Cys Phe Arg Asn Val Gln Asp Asn Cys Cys Leu 325 330 Arg Pro Leu Tyr Ile Asp Phe Lys Arg Asp Leu Gly Trp Lys Trp Ile 340 345 350 His Glu Pro Lys Gly Tyr Asn Ala Asn Phe Cys Ala Gly Ala Cys Pro Tyr Leu Trp Ser Ser Asp Thr Gln His Thr Lys Val Leu Ser Leu Tyr 370 375 Asn Thr Ile Asn Pro Glu Ala Ser Ala Ser Pro Cys Cys Val Ser Gln 385 390 395 Asp Leu Glu Pro Leu Thr Ile Leu Tyr Tyr Ile Gly Asn Thr Pro Lys 405 410 Ile Glu Gln Leu Ser Asn Met Ile Val Lys Ser Cys Lys Cys Ser 420 425 <210> 28 <211> 1272 <212> DNA <213> Artificial Sequence <220> <223> Fusion oligonucleotide <220> <221> CDS <222> (1)..(1272)

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tac Tyr 305	aag Lys	gat Asp	gac Asp	gac Asp	gac Asp 310	aag Lys	gcc Ala	ctg Leu	gac Asp	acc Thr 315	aat Asn	tac Tyr	tgc Cys	ttc Phe	cgc Arg 320	960
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Leu Aro	g Leu	Thr	Ser	Pro	Pro 55	Glu	Pro	Ser	Val	Met 60	Thr	His	Val	Pro	

Tyr Gl
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Pro Gly Gln Gly Ser Gln Arg Lys Lys Arg Ala Leu Asp Thr Asn Asp 290 295 300

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		gtg Val														768
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		ttc Phe														1008
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		tat Tyr 355	_			_								_	_	1104
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1284

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- Tyr Gln Val Leu Ala Leu Tyr Asn Ser Thr Arg Glu Leu Leu Glu Glu 65 70 75 80
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His	Glu	Val	Met	Glu 245	Ile	Lys	Phe	Lys	Gly 250	Val	Asp	Asn	Glu	Asp 255	Asp
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Pro	Gly 290	Gln	Gly	Ser	Gln	Arg 295	Lys	Lys	Arg	Ala	Leu 300	Asp	Thr	Asn	Ser
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	tg cta al Leu														385
	cc ccc nr Pro														433
	cg gtg la Val														481
	gg ctc rg Leu 160														529
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1353

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Lys	Leu	Lys	Val	Glu 165	Gln	His	Val	Glu	Leu 170	Tyr	Gln	Lys	Tyr	Ser 175	Asn
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Ser	Gly	Arg	Arg	Gly 245	Asp	Leu	Ala	Thr	Ile 250	His	Gly	Met	Asn	Arg 255	Pro
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ctg t Leu S 30																145
atc (193
ccc (Pro I																241
ctg (Leu <i>l</i>	_				_		_	_		_	_		_	_	-	289
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cgc o Arg V 110																385
ggc a Gly :																433
gaa q Glu <i>I</i>																481
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- Gln Gly Asp Val Pro Pro Gly Pro Leu Pro Glu Ala Val Leu Ala Leu 65 70 75 80
- Tyr Asn Ser Thr Arg Asp Arg Val Ala Gly Glu Ser Val Glu Pro Glu 85 90 95
- Pro Glu Pro Glu Ala Asp Tyr Tyr Ala Lys Glu Val Thr Arg Val Leu 100 105 110
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115 120 125

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